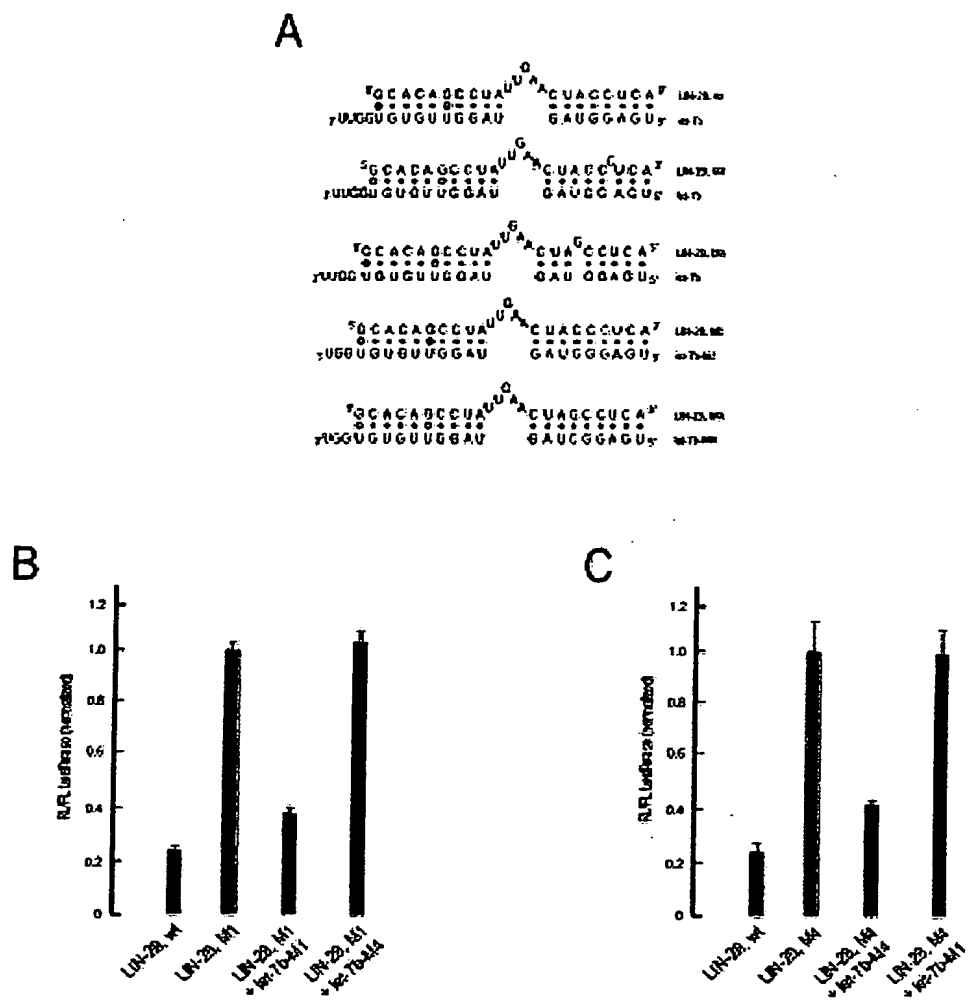


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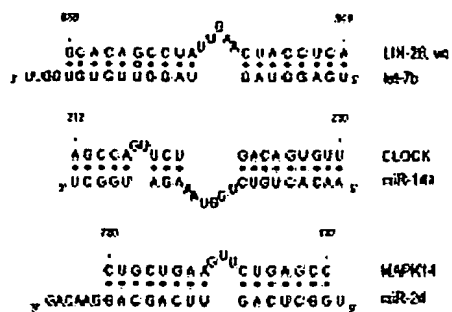
Figure 2



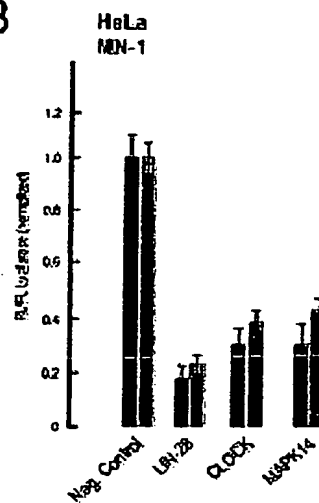
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Figure 3

A



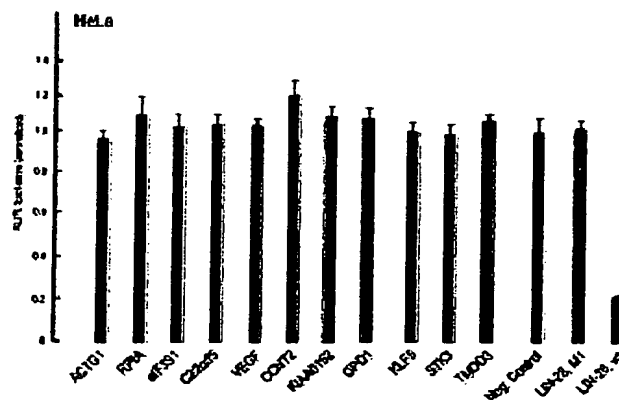
B



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Figure 4

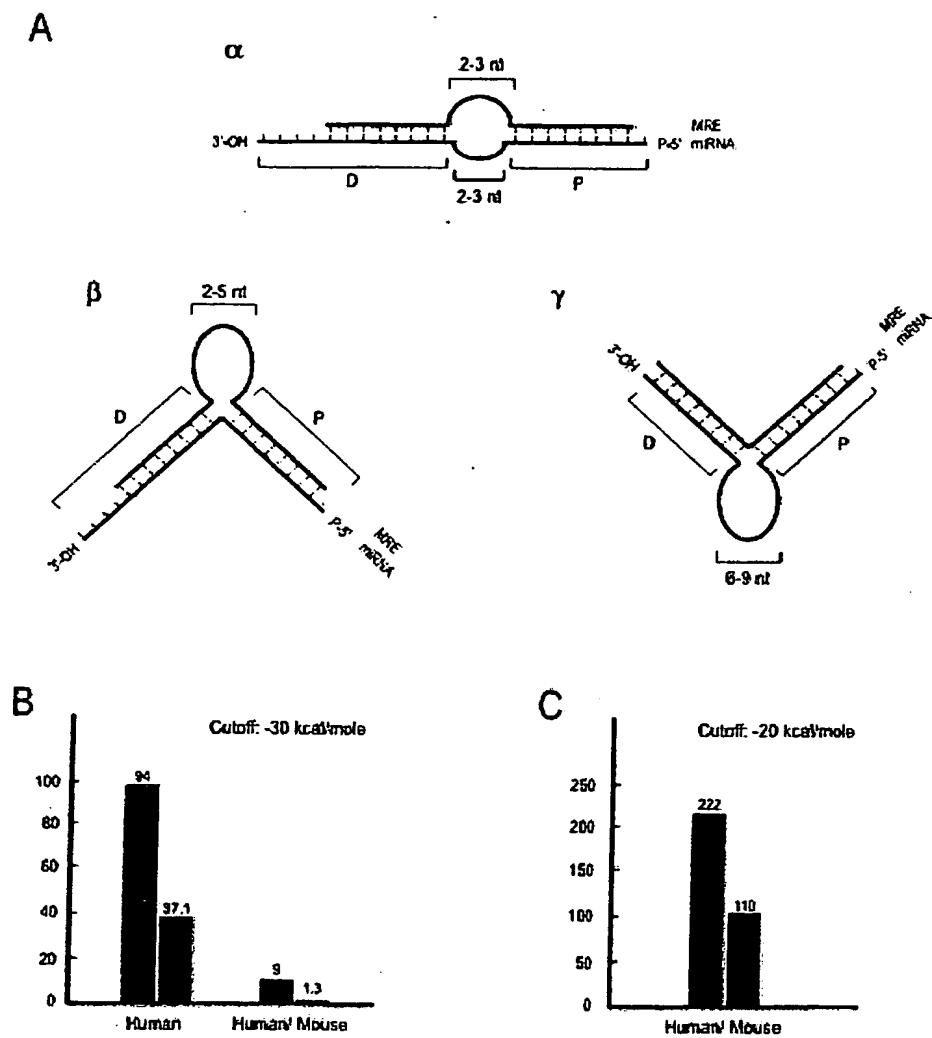
miRNA	Target Annotation	Hit location (miRNA 3' end)	miRNA Binding Site (Top Strng: Target; Bottom Strng: miRNA)
let-7b	Actin gamma 1 (ACTG1; NM_001614)	1294	5' ---cagaca-cctccatgctagccca--- 3' 3' ---gtgtctcga---tcat-ggagc--- 5'
let-7b	Ribose 5-phosphate isomerase A (RPIA; NM_144563)	1013	5' -----cacagccaggtggagcaccac--- 3' 3' ---gggtgtgctgg-----atg-atggag--- 5'
let-7c	Eucaryotic translation initiation factor 3, subunit 1 alpha (eIF3S1; NM_003758)	1140	5' ---acacagcccttaccctaccac--- 3' 3' ---tatgttggg---gg-atggag--- 5'
miR-15	Chromosome 22 open reading frame 5 (C22orf5; NM_012264)	3152	5' -----cagggca---atgtgctgct--- 3' 3' ---gtgttctgtataccagcaga--- 5'
miR-16	Vascular endothelial growth factor (VEGF; NM_003376)	1560	5' ---cgcacatcttcttctctctgctgcca--- 3' 3' ---gggtt---tatata-tgacagcagct--- 5'
miR-16	Cyclin T2 (CCNT2), transcript variant a (NM_001241)	2254	5' -----caatccctcgaagctgctgct--- 3' 3' -----gggttatata-tgacagcaga--- 5'
miR-24	KIAA0152 gene product (NM_014730)	4671	5' -----cctcgtc---cctgagcca--- 3' 3' ---gacagggacgactctgcccggc--- 5'
miR-103	Glycerol-3-phosphate dehydrogenase 1 (soluble) (GPD1; NM_005276)	1368	5' ---tca-agccc---cag-gccgc--- 3' 3' ---agttctgggacatgctccgagc--- 5'
miR-141 (miR-157)	Kruppel-like factor 5 (intestinal) (KLF5; NM_001730)	2033	5' -----caatcatttactgacagctgct--- 3' 3' -----tcggtag-aaat-ggtctgcaaaa--- 5'
miR-141 (miR-157)	Serine/threonine kinase 3 (STE20 homolog, yeast; STK3; NM_006281)	2756	5' ---ggaatctt---gacagctgtc--- 3' 3' ---cgttagaactggcctgctcaga--- 5'
miR-145	Tropomodulin 3 (ubiquitous) (TMOD3; NM_014547)	1225	5' ---tccctgggtgaaagggaaggaacgga--- 3' 3' ---aggaacct---cttcgacct--- 5'



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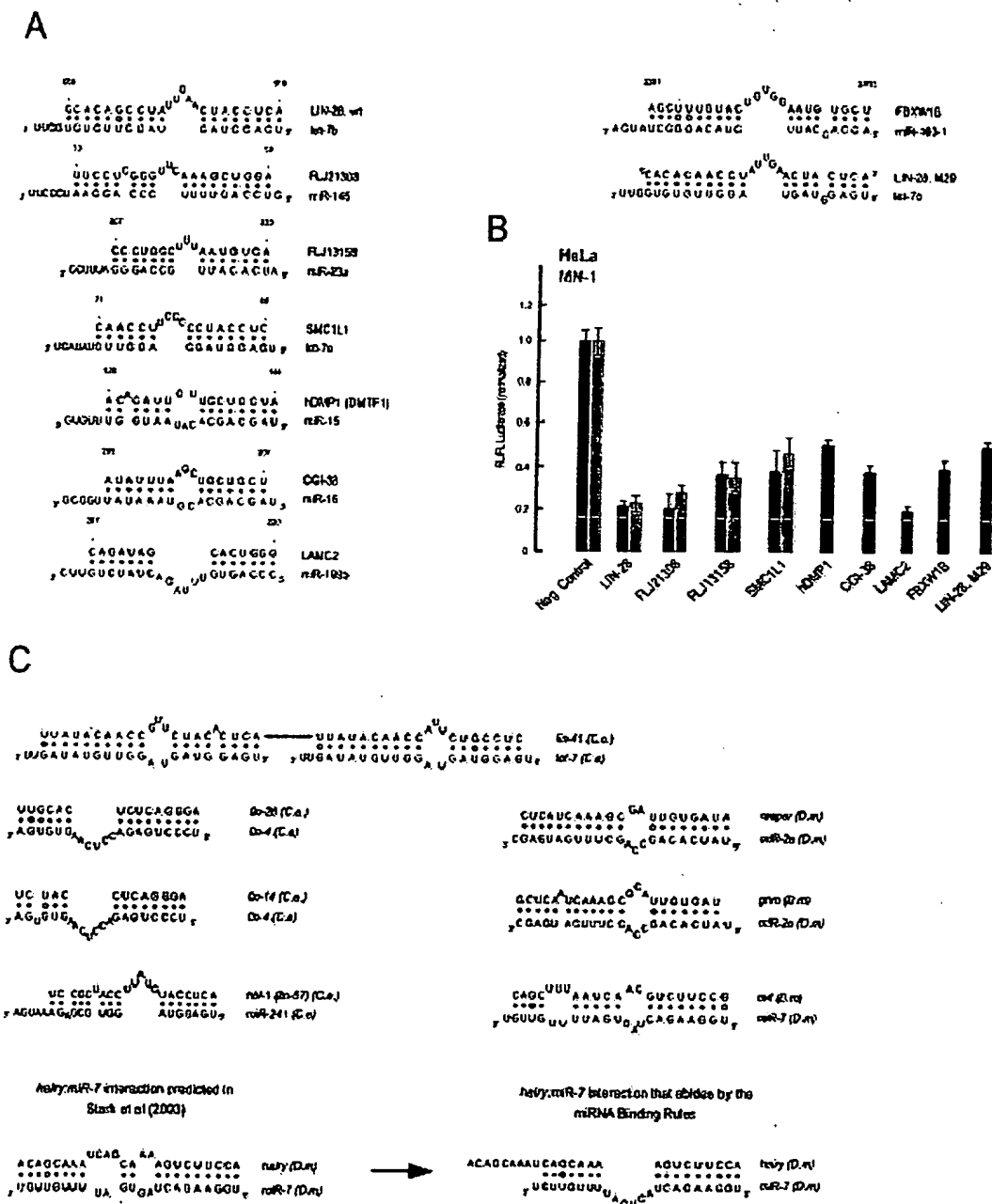
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Figure 5.



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Figure 6



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